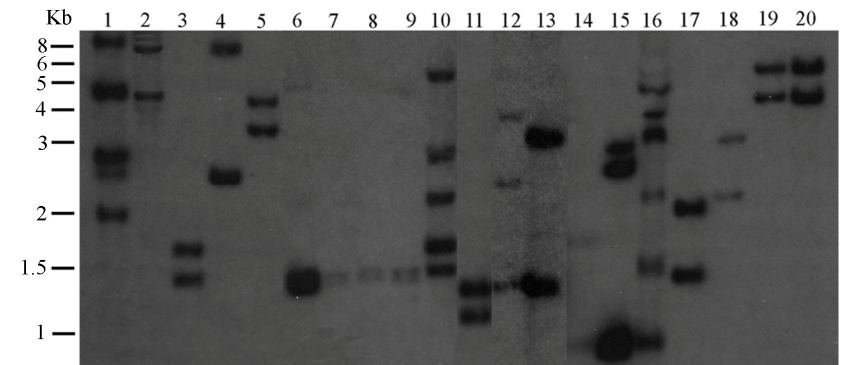
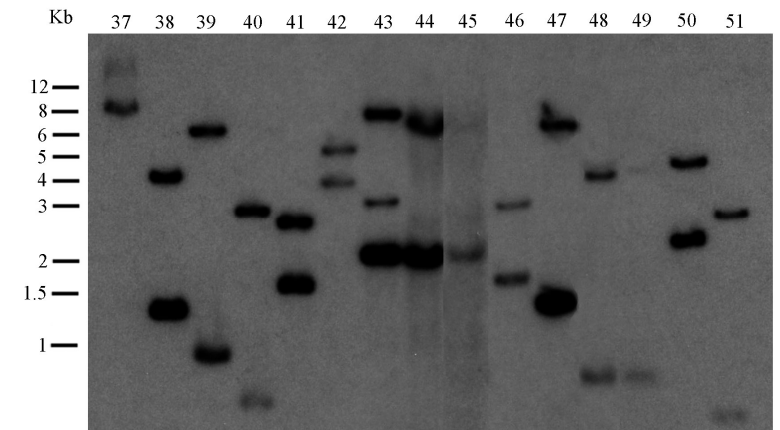
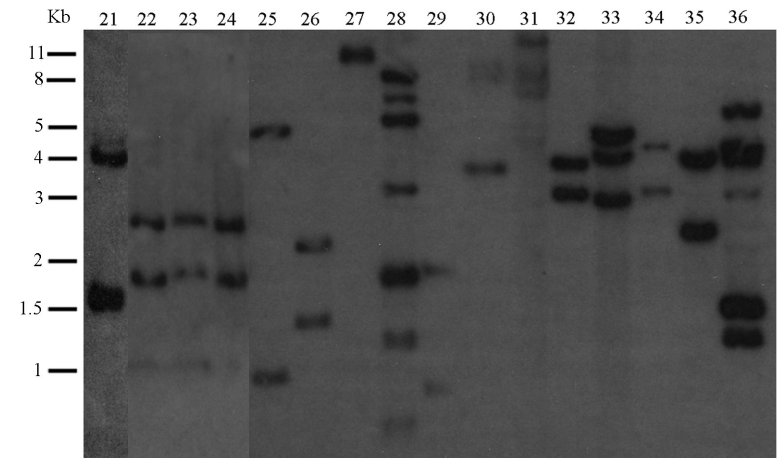


SOUTHERN 1				
Lane	Mutants	Size (bp)	Size (bp)	Comments
1 - 2	<i>bot1</i> Δ	4890	8645	Obtained the bands of the expected size in well #2. This mutant was used for further experiments
3	<i>byr4</i> Δ	1599	1557	Obtained the bands of the expected size
4	<i>bur6</i> Δ	2622	8260	Obtained the bands of the expected size
5	<i>cdc14</i> Δ	3740	4835	Obtained the bands of the expected size
6 - 10	<i>cet1</i> Δ	3382	1353	Only the lower band is correct in # 6 - 9. The mutant corresponding to well #6 was used for further experiments
11	<i>erg8</i> Δ	1264	1049	Obtained the bands of the expected size
12 - 13	<i>fba1</i> Δ	8192	1266	Mutant corresponding to well #13 has two hybridization bands, although only the lower is of the expected size. This mutant was used for further experiments
14	<i>fol1</i> Δ	1720	849	Obtained the bands of the expected size
15 - 16	<i>hrb1</i> Δ	3293	816	Obtained the bands of the expected size in #15 where there is also one additional hybridization band. The mutant corresponding to well #15 was used for further experiments
17	<i>kei1</i> Δ	1316	2051	Obtained the bands of the expected size
18	<i>ktr3</i> Δ	3682	2115	Obtained the bands of the expected size
19 - 20	<i>mdm10</i> Δ	5999	4572	Obtained the bands of the expected size in both candidate transformants, and the one corresponding to well #19 was used for further experiments

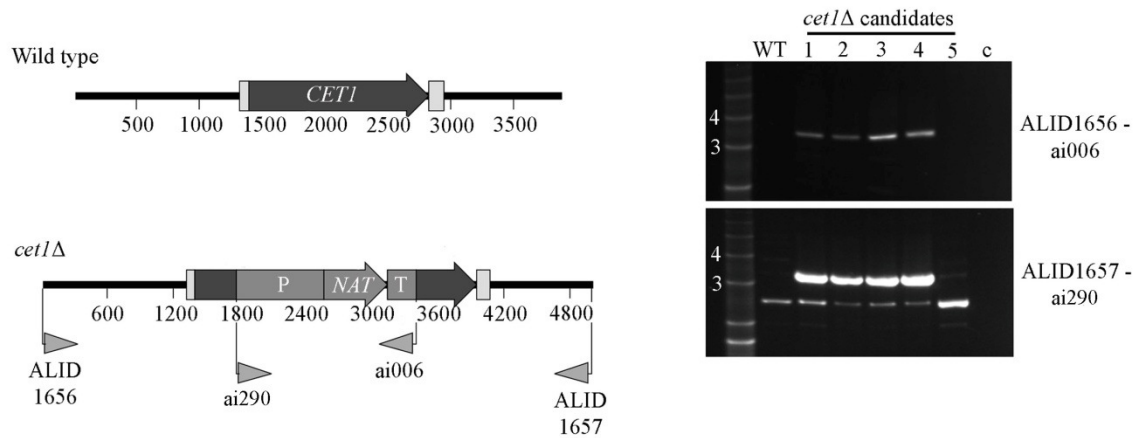


SOUTHERN 2				
Lane	Mutants	Size (bp)	Size (bp)	Comments
21	<i>mdm34Δ</i>	1605	4737	Obtained the bands of the expected size
22 - 24	<i>mgm101Δ</i>	2617	1880	Obtained the bands of the expected size in all the three <i>mgm101Δ</i> candidate mutants selected. The mutant corresponding to well #23 was used for further experiments
25	<i>mmm1Δ</i>	4919	999	Obtained the bands of the expected size
26	<i>mrpl7Δ</i>	2146	1381	Obtained the bands of the expected size
27	<i>mrpl31Δ</i>	11207	12706	Obtained the bands of the expected size
28	<i>mrps18Δ</i>	6672	17142	There are multiple hybridization bands. This mutant was used for further experiments
29	<i>mvd1Δ</i>	2096	883	Obtained the bands of the expected size
30 - 31	<i>nam9Δ</i>	7856	9211	Obtained the bands of the expected size in #31. This mutant was used for further experiments
32	<i>pwp1Δ</i>	2933	3674	Obtained the bands of the expected size
33- 34	<i>cdc1Δ</i>	2571	4039	Both mutants do not have the bands of expected size, even though the PCR is positive (see below). The mutant in well #33 was used both for genetic analysis and haploinsufficiency, while the mutant in well #34 was only used for the haploinsufficiency experiment as it is unable to produce spores for genetic analysis
35	<i>rib2Δ</i>	3867	2289	Obtained the bands of the expected size
36	<i>rib3Δ</i>	2292	3768	There are multiple hybridization bands. This mutant was used for further experiments

SOUTHERN 3				
Lane	Mutants	Size (bp)	Size (bp)	Comments
37	<i>rsa4Δ</i>	15235	8523	Obtained the bands of the expected size
38	<i>rsc9Δ</i>	1382	4221	Obtained the bands of the expected size
39	<i>rsm18Δ</i>	6730	945	Obtained the bands of the expected size
40	<i>saf2Δ</i>	2995	592	Obtained the bands of the expected size
41	<i>sec5Δ</i>	1660	2743	Obtained the bands of the expected size
42	<i>sen54Δ</i>	3877	5331	Obtained the bands of the expected size
43 - 45	<i>sfi1Δ</i>	2103	7811	Obtained the bands of the expected size in #44 and #45; the mutant corresponding to well #44 was used for further experiments
46	<i>CNAG_00592Δ</i>	1713	3036	Obtained the bands of the expected size
47	<i>thp1Δ</i>	5456	1451	Obtained the bands of the expected size
48 - 49	<i>tim54Δ</i>	3901	279	Same hybridization pattern for both <i>tim54Δ</i> candidates; only the higher band is of the expected size. Both mutants were used for further experiments.
50	<i>trl1Δ</i>	4390	2277	Obtained the bands of the expected size
51	<i>trr1Δ</i>	2703	564	Obtained the bands of the expected size

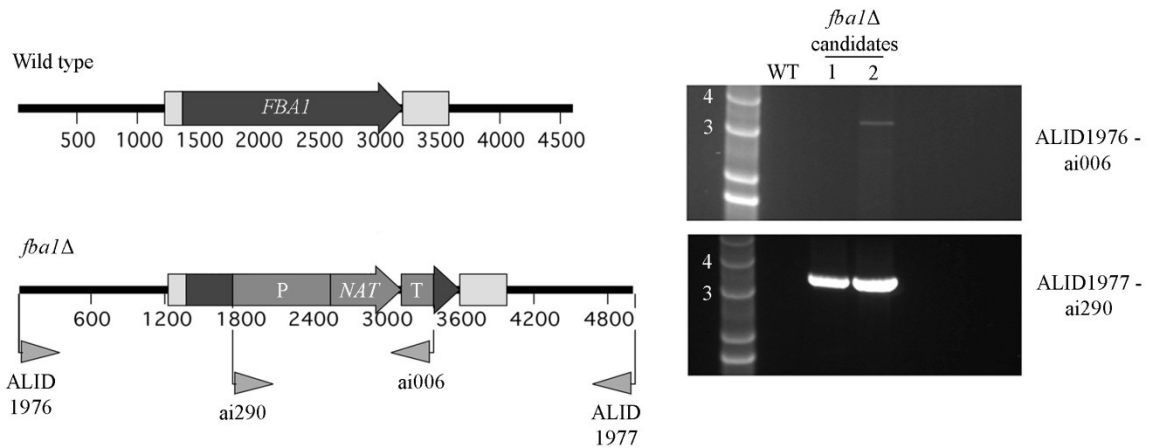


A) Gene replacement and PCR confirmation for the gene *CET1*



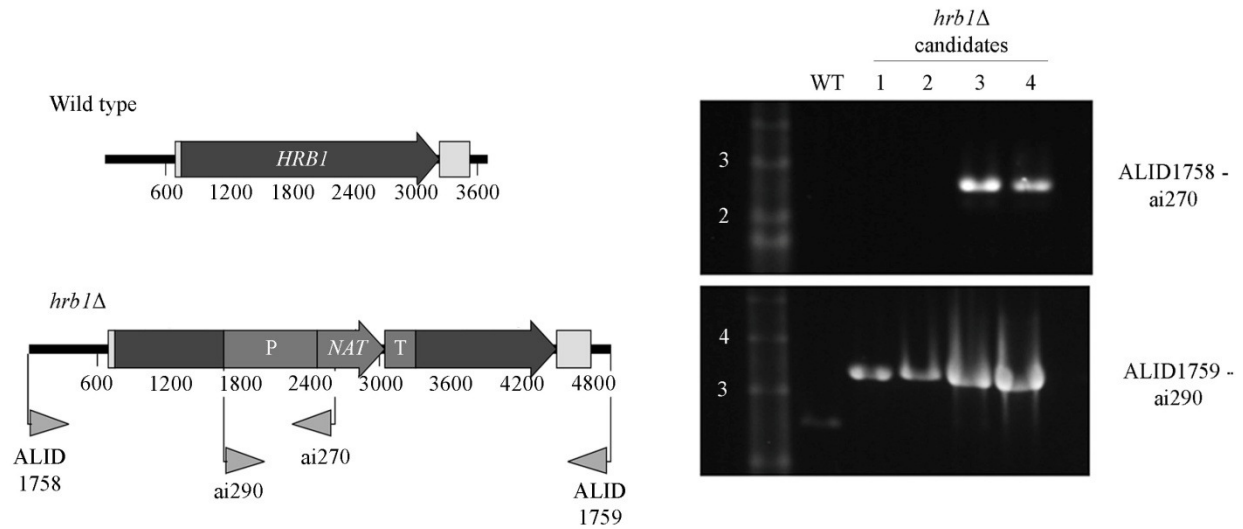
Comment: *cet1Δ* candidates #1 - 5 correspond to wells #6 - 10 of the Southern blot 1. Candidates *cet1Δ* #1, #2, #3 and #4 have the same amplification pattern, and *cet1Δ* #1 (#6 of Southern 1) was used for further experiments. PCR with primers ALID1657-ai290 produced a non-specific amplicon of ~2.5 Kb. c: water control.

B) Gene replacement and PCR confirmation for the gene *FBA1*



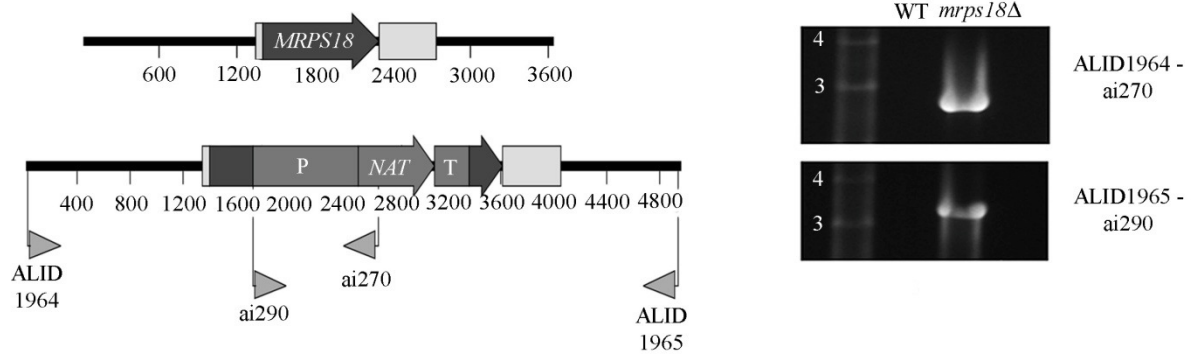
Comment: *fba1Δ* candidates #1 and #2 correspond to wells #12 and #13 of the Southern blot 1. PCR for candidates *fba1Δ* #2 showed correct integration of the gene replacement construct, while in candidate *fba1Δ* #1 most likely only the right side is correctly integrated. Therefore, mutant *fba1Δ* #2 (#13 of Southern 1) was used for further experiments.

C) Gene replacement and PCR confirmation for the gene *HRB1*



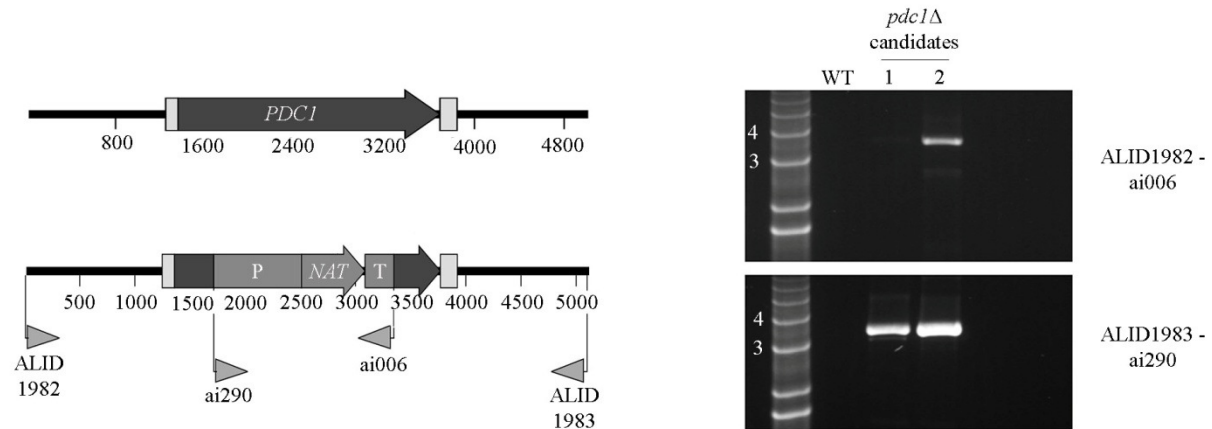
Comment: PCR for candidates *hrb1Δ* #1 and #2 shows that probably only the right side of the gene replacement construct is correctly integrated, while for candidates *hrb1Δ* #3 and #4 both left and right sides are correctly integrated. *hrb1Δ* candidates #3 and #4 correspond to well #15 and #16 of the Southern blot 1. Mutant *hrb1Δ* #3 (#15 of Southern 1) was used for further experiments.

D) Gene replacement and PCR confirmation for the gene *MRPS18*



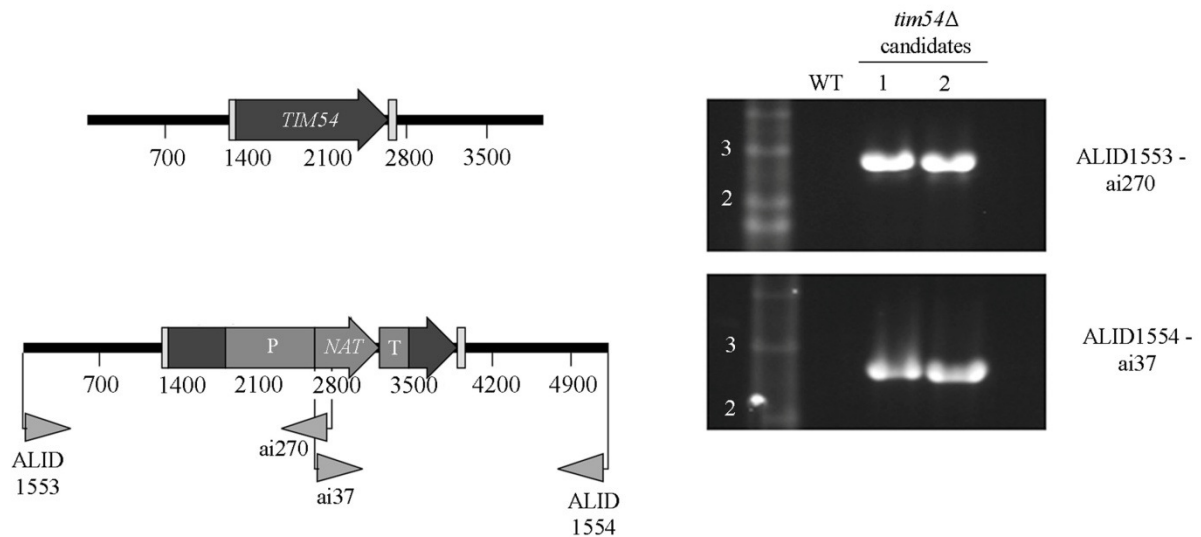
Comment: one candidate *mrps18Δ* strain was isolated (#28 of Southern 2). The PCR produced bands of the expected size, and this mutant was used for further experiments.

E) Gene replacement and PCR confirmation for the gene *PDC1*



Comment: *pdclΔ* candidates #1 and #2 correspond to wells #33 and #34 of the Southern blot 2; PCR showed correct integration of the gene replacement construct, even though for candidate *pdclΔ*#1 the amplicon obtained using primers ALID1982-ai006 was weak. The mutant *pdclΔ*#1 was used both for genetic analysis and haploinsufficiency, while the mutant *pdclΔ*#2 was only used for the haploinsufficiency experiments as the strain was unable to produce spores for genetic analysis.

F) Gene replacement and PCR confirmation for the gene *TIM54*



Comment: *tim54Δ* candidates #1 and #2 correspond to wells #48 and #49 of the Southern blot 3; PCR showed correct integration of the gene replacement construct for both candidates, which were both used for further experiments.